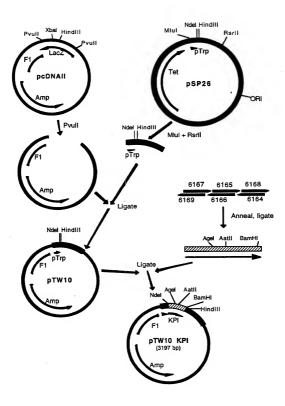
Title: PROTEASE INHIBITOR PEPTIDES
Inventor(s): R. Tyler WHITE et al.
Appl. No.: 09/201,715

Figure 1



Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201.715

Figure 2

Ndel

TATG AAA CAA AGC ACT ATT GCA CTG GCA CTC TTA CCG TTA CTG TTT ACC CTT GTG ACA AAA AC TTT GTT TCG TGA TAA CCT GAC CGT GAG AAT GGC AAT GAC AAA TGG GGA CAC TGT TTP. Met Lys Gin Ser Thr ile Ala Leu Ala Leu Leu Pro Leu Leu Phe Thr Pro Val Thr Lys

(PI Agel

OCC GAG OTO TOO TOT GAA CAA GOT GAG ACC GOT GOO TOO GOT GOA ATG ATC TOC GOC TGG GOTTO CAC GAGA CITT OTT GOA CIT TGG CCA GOG AGG GCA GOT TAC TGG AGG GCG ACC Ala Glu Vai Cys Ser Glu Glin Ala Glu Thr Gly Pro Cys Arg Ala Met Ile Ser Arg Trp

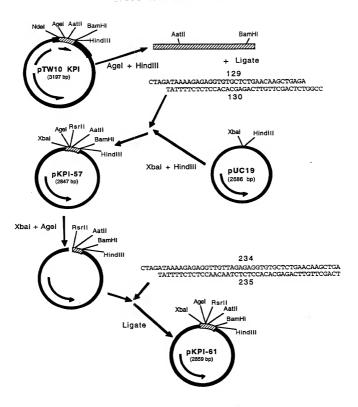
Aatil

TAC TIT GAC GIC ACT GAA GGT ANG TGC GCT CCA TIC TIT TAC GGC GGT TGC GGC GGC ANG ANG AAA CIT GCM TGA CIT CCA TIC AGG GGA GGT ANG AAA ATG CCG CCA GG CGC GGC TG Flyr Phe Asp Val Thr Giu Giy Lys Cys Ala Pro Phe Phe Tyr Gly Giy Cys Giy Gly Asn

BamHI Hir

CGT AAC AAC TIT GAC ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TTA GAC ATT TTG AAA CTG TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT ACG CGA TAA ATT CGA PATG ASA Ash Phe Asp Thr Glu Glu Tyr Cys Met Ala Val Cys Gly Ser Ala Ile

Appl. No.: 09/201,715



Title: PROTEASE INHIBITOR

PEPTIDES Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 4

Xbal

Rsrii

KPI (1-57) Agel

CTA GAT AAA AGA GAG GTG TGC TCT GAA CAA GCT GAG ACC GGT CCG TGC CGT TA TTT TCT CTC CAC ACG AGA CTT GTT CGA CTC TGG CCA GGC ACG GCA Leu Asp Lys Arg Glu Val Cys Ser Glu Gin Ala Glu Thr Gly Pro Cys Arg

Aatii

GCA ATG ATC TCC CGC TGG TAC TTT GAC GTC ACT GAA GGT AAG TGC GCT CCA
CGT TAC TAG AGG GCG ACC ATG AAA CTG CAG TGA CTT CCA TTC ACG CGA GGT Ala Met Ile Ser Arg Trp Tyr Phe Asp Val Thr Giu Giy Lys Cys Ala Pro

TTC TTT TAC GGC GGT TGC GGC GGC AAC CGT AAC AAC TTT GAC ACT GAA GAG AAG AAA ATG CCG CCA ACG CCG CCG TTG GCA TTG TTG AAA CTG TGA CTT CTC

Phe Phe Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp Thr Glu Glu

> BamHI HindIII

TAC TOC ATG GCA GTG TGC GGA TCC GCT ATT TA ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA Tyr Cys Met Ala Val Cys Gly Ser Ala lie

Figure 5

Kbal KPI (-4-57)

Agel

CTA GAT ANA AGA GAT OFF AGA GAG GTG TOC TCT GAA CAA GTT GAG AGA GAC GAT

TA TTT TCT CTC CAA CAA TCT CTC CAC AGG AGA CTT GTT GGA CTC TOG CCA

**Leu Asp Lys Arg Glu Vai Vai Arg Glu Vai Cys Ser Glu Gin Ala Glu Thr Gly

AatII

CCG TGC CGT GCA ATG ATC TCC CGC TGG TAC TTT GAC GTC ACT GAA GGT AAG TGC GGC AGC GCA GCT TAC TAG AGG GCG ACC ATG AAA CTT CAC TTC ACT PPro Cys Arg Ala Met lie Ser Arg Trp Tyr Phe Asp Val Thr Glu Gly Lys Cys

OCT CCA THE THE TAC GGC GGT TOC GGC GGC AAC CGT AAC AAC THE GAC ACT GAA CGA GGT AAG AAA ATG CCG CCA ACG CCG CCG THE GCA THE THE AAA CHE TAG CHE AAA AP THE CHE CAA CCC CCG CCG THE GCA THE THE THE AAA CHE TAG CHE AAA AP THE AAF THE GIU

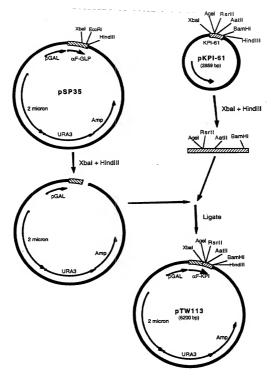
BamHI HindIII

GAG TAC TCC ATG GCA GTG TCC GCA ATT TA

CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA

> GIU Tyr Cys Met Ala Val Cys Gly Ser Ala IIe

Figure 6



Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 7

Y -	fа	C	tο	r	

ATG AGA TTT CCT TCA ATT TTT ACT GCA GTT TTA TTC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA Met Arg Phe Pro Ser lie Phe Thr Ala Val Leu Phe Ala Ala Ser Ser Ala Leu Ala

GCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CCA GGT CAG TTG TGA TT GTC ACT TCA CTT TCC CGT GTT TAA GGC CAC CTT CGA CAG ALIA VAI AND VAI AND THE THE THE GLU ANN GU THE ALIA GTO ALIA GAI ALIA VAI

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC
TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTG Ille Civ Tyr Leu Asp Leu Giu Giv Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn

AGC ACA AAT AAC GOG TTA TTG TTT ATA AAT ACT ACT ATT GCC AGC ATT GCT GCT AAA TCG TGT TTA TTG CCC AAT AAC AAA TAT TTA TGA TGA TAA CGG TCG TAA CGA CGA TTT ▶ Ser Thr Asn Asn Gly Leu Leu Phe IIe Asn Thr Thr IIe Ala Ser IIe Ala Ala Lys

KPI(-4-57) Xbal

GAA GAA GGG GTA TCT CTA GAT AAA AGA GAG GTT GTT AGA GAG GTG TC TCT GAA CAA TCT CTC CCC AAT AGA GAT CTA TCT TCT CTC CAA CAA TCT CTC CAC AGA GAG ACT GTT CGU GU GU YAI Ser Leu Asp Lys Arg G u Vai Vai Arg Gu Vai Cys Ser Giu Gin

Rsrll

Agel Aatli

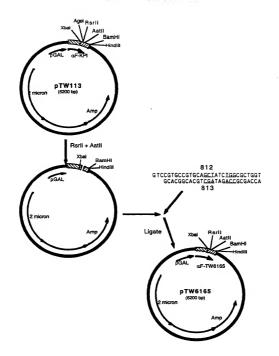
GCT GAG ACC GGT CCG TGC CGT GCA ATG ATC TCC CGC TGG TAC TTT GAC GTC ACT GAA CGA CTC TGG CCA GCC ACG GCA CGT TAC TAG AGG GCG ACC ATG AAA CTG CAG TGA CTT PALA GLU Thr GLV Pro Cvs Arg Ala Met II e Ser Arg Trp Tvr Phe Asp Val Thr Glu

BamHI

ACT GAA GAG TAC TOC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A Thr Glu Glu Tyr Cys Met Ala Val Cys Gly Ser Ala Ile

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 9



Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201 715

Figure 10

pTW 6165

~-	fа	CI	'n	1

ATG AGA TIT CCT TCA ATT TIT ACT GCA GTT TTA TTC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA Met Arg Phe Pro Ser ile Phe Thr Ala Vai Leu Phe Ala Ala Ser Ser Ala Leu Ala OCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CCA GGT CAG TTG TGA TGT TGT CTT CTA CTT TGC CGT GTT TAA GGC CGA CTT CGA CAG ÀAIa Pro Vai Asa Tht Thr Thr Giu Asp Giu Thr Ala Gin Ile Pro Ala Giu Ala Vai ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTG File Gly Tyr Leu Asp Leu Glu Gly Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn KPI(-4-57: M15A, S17W) Xbal

GAA GAA GOG GTA TOT CTA GAT AAA AGA GAG GTT GTT AGA GAG GTG TOC TOT GAA CAA CTT CTT CCC AAT AGA GAT CTA TTT TCT TCT CAC CAA TCT CTC CAC AGG AGA CTT GTT GTU GTU GTY VAI Ser Lou Asp Lys Arg GTU VAI VAI Arg GTU VAI Cys Ser GTU GTN

Agel GCT GAG ACC GGT CCG TGC CGT GCA GCT ATC TGG CGC TGG TAC TTT GAC GTC ACT GAA
CGA CTC TGG CCA GGC ACG GCA CGT CGA TAG ACC GCG ACC ATG AAA CTG CAG TGA CTT Ala Giu Thr Giv Pro Cvs Arg Ala Ala IIe Trp Arg Trp Tvr Phe Asp Val Thr Giu GGT AAG TGC GCT CCA TTC TTT TAC GGC GGT TGC GGC GGC AAC CGT AAC AAC TTT GAC CCA TTC ACG CGA GGT AAG AAA ATG CGG CCA ACG CGG CGG TTG GGC TTG GGC TTG TTG AAA CTG PG YL Lys Cys Ala Pro Phe Phe Tyr Gly Gly Cys Gly Asn Arg Asn Asn Phe Asp

ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC AGG CCT AGG CGA TAA ATT CGA A FAR CTI GU GIU Tyr Cys Met Ala Val Cys Gly Ser Ala IIe

812	
GTCCGTGCCGTGCACCLATCLGGCGCTGGTACTTTGACGT GCACGGCACGTLGATAGALCGCGACCATGAAAC 813	pTW6165 KPI(-4-57; M15A, S17F)
814 GTCCGTGCCGTGCAGCTATCTACCGCTGGTACTTTGACGT GCACGGCACG	pTW6166 KPI(-4-57; M15A, S17Y)
867 GTCCGTGCGTGCAIIGATCIICCGCTGGTACTTTGACGT GCACGGCACGTAACTAGAAGGCGACCATGAAAC 868	pTW6175 KPI(-4-57; M15L, S17F)
1493 GTCCGTGCCGTGCALIGATCIACCGCTGGTACTTTGACGT GCACGGCACGTAACTAGATGGCGACCATGAAAC 1494	pBG028 KPI(-4-57; M15L, S17Y)
925 GTCCGTGCCGTGCAATGCACTICCGCTGGTACTTTGACGT GCACGGCACGTTACGTGAAGGCGACCATGAAAC 926	pTW6183 KPI(-4-57; I16H, S17F)
927 GTCCGTGCCGTGCAATG <u>CACTAC</u> CGCTGGTACTTTGACGT GCACGGCACGTTAC <u>GTGATG</u> CGACCATGAAAC 928	pTW6184 KPI(-4-57; I16H, S17Y)
929 GTCCGTGCCGTGCAATG <u>CACLGG</u> CGCTGGTACTTTGACGT GCACGGCACGTTAC <u>GTGACC</u> GCGACCATGAAAC 930	pTW6185 KPI(-4-57; I16H, S17W)
863 GTCCGTGCCGTGCAGCICACTCCCGCTGGTACTTTGACGT GCACGGCACGTCGAGTGAGGCGACCATGAAAC 864	pTW6173 KPI(-4-57; M15A, I16H)
865 GTCCGTGCCGTGCAITGCACTCCCGCTGGTACTTTGACGT GCACGGCACGTAACGTA	pTW6174 KPI(-4-57; M15L, I16H)

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201.715

Figure 12

pTW 6166

α-factor

ANG AGA TIT OUT TOA AIT TIT ACT GOA GIT THA THE GCA GCA TOC TOC GCA THA GCT TAC TOT ANA GGA AGT TAA ANA TGA COT CAA ANT ANG COT COT AGG AGG COT ANT CGA PMet Arg Phe Pro Ser IIe Phe Thr AIa Val Leu Phe AIa Aia Ser Ser Aia Leu Aia

CCT CCA OTC AND ACT ACA ACA GAA GAR GAA ACG CCA ACA ATT CCG CCT GAA CCT CCA CCA GCT CAG TTG TCA TCT TCT CTT CTA CTT TCC CCT TTA GCC CCA CTT CAA CCT CCA CAG PAIa Pro Val Aan Thr Thr Thr Giu Asp Giu Thr Aia Gin I is Pro Aia Giu Aia Vai

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC
TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTA
Pile Giy Tyr Leu Aap Leu Giu Giy Aap Phe Asp Vai Ala Vai Leu Pro Phe Ser Asn

ACC ACA AAT AAC GGG TTA TTG TTT ATA AAT ACT ACT ACT ATT GCC ACC ATT GCT GCT AAA TG TGT TTA TTG CCC AAT AAC AAA TAT TTA TGA TGA TGA TGG TGG TAA CAG GGA TTT Ser Tha An An GIG Leu Leu Phe lie Asn Thr Thr Ile Ala Ser lie Ala Ala Lys

al KPI(-4-57; M15A, S17Y)

GAA GAA GGG GTA TOT CTA GAT AAA AGA GAG GTT GTT AGA GAG GTG TGC TCT GAA CAA CTT GTT CTC CCC CAT AGA GAT CTA TTT TCT CTC CAC CAC CAC CAC CTC GTT GTU GIU GIU GIV VAI Sor I GUU Asp Lys Arg Giu Vai Vai Arg Giu Vai Cys Sor Giu Gin

RsrII

Agel

OUT GAG ACC GOT CCG TGC AGT ACC TAC CGC TGG TAC TIT GAC GTC ACT GAA

CGA CTC TGG CGA GGC ACG GGA AGG TGGA TAC ATG GAC ACC ATG AAA CTG CAG TGA CTT

Ala Glu Thr Gly Pro Cys Arg Ala Ala II e Tyr Arg Trp Tyr Phe Asp Val Thr Glu

GOT AAG TOC GOT COA TITC TITT TAC GOC GOT TOC GOC GOC AAC COT AAC AAC TITT GAC COA TITC ACG GOT AAG AAA ATG COG COA ACG GOC GOG TITG GOA TITG TITG AAA CTG PGIJ Lys Cys Ala Pro Phe Phe Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp

BamHI HindIII

ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A PThr Glu Glu Tyr Cys Mat Ala Val Cys Gly Ser Ala 11e Title: PROTEASE INHIBITOR
PEPTIDES
Inventor(s): R. Tyler WHITE et al.

Appl. No.: 09/201,715

Figure 13

pTW 6175

α-1	ıa	C	ta	۳

Ser Thr Ash Ash Gly Leu Leu Phe IIe Ash Thr Thr IIe Ala Ser IIe Ala Ala Lys KPI(-4-57; M15L, S17F)

RsrII

BamHI HindIII
ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T
TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A
Thr Gu Giu Tyr Cys Mat Ala Val Cys Giy Sar Ala IIe

Title: PROTEASE INHIBITOR PEPTIDES Inventor(s): R. Tyler WHITE et al.

Appl. No.: 09/201.715

Figure 14

pBG028

α-factor

ATTS AGA TITT COT TOA ATTT TITT ACT GOA GITT TITA TITO GOA GOA TOO TOO GOA ATTA GOT TAC TOT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA PMet Arg Phe Pro Ser iie Phe Thr Aia Vai Leu Phe Aia Aia Ser Ser Aia Leu Aia

GCT CCA GTC AND ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC GAG GGT CAG GTC TCA GTT TCT CTT CTA CTT TCC GCT GTT TTA GGC GCA CTT CGA CAG $^{\circ}$ Ala Pro Val Asa Thr Thr Glu Asp Glu Thr Ala Gin lie Pro Ala Glu Ala Val

ATC GOT TAC THA GAT THA GAD GOG GAT THE GAT GTT GCT GTT THE CCA THT TCC AAC THA GCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CAA CAA AAC GTT AAA AAG TTA FILE GBY Tyr Leu Aay Leu Gly Aap Fhe Aap Val Ala Val Leu Pro Fhe Sar Aan

AGC ACA ANT ANG GGG THA THG THT ATM ANT ACT ACT NIT GCC AGC ATT GCT GCT ANA TGG TGT THA THG CCC ANT ANG ANA TAT THA TGA TGA TAA TAA CGG TGG TAA CGA GGA THT > Ser Thr Asn Asn Gly Leu Leu Phe II e Asn Thr Thr II e Ala Ser II e Ala Ala Lys

hal KPI(-4-57; M15L, S17Y)

GAA GAA GGG GTA TOT CTA GAT AAA AGA GAG GTT GTT AGA GAG GTG TGC TGA CAA CTT CTT CCC CAT AGA GAT CTA TTT TCT CTC CAA CAA TCT CTC CAC ACG AGA CTT GTT GTU GTU GTU VAI SAT LGU ASP LYS AT GT GTU VAI A TG GTU VAI CyS SAT GTU GT

Rsril

Agel Agt

GOT AND TOC GCT CCA TTC TITT TAC GGC GGT TCC GGC GGC AAC GCT AAC AAC TTT GAC CCA TCC ATC ACG CGA ACG CGC ATC GCC ATC TTT DAAC CTG FCI LY Cys Cys Ala Pro Phe Phe Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp

BamHl HindIII

ACT GAA GAG TAC TOC ATG GCA GTG TOC GCA TCC GCT ATT TAA GCT T TCA CTT CTC ATG AGG TAC COT CAC ACG CCT AGG CCA TAA ATT CGA A TTR Glu Glu Tyr Cys Met Ala Val Cys Gly Ser Ala lie

Figure 15

pTW6183

α-factor

ATG AGA TITT ÉCT TCA AIT TITT ACT GCA GTT TTA TTC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA PMet Arg Phe Pro Ser IIe Phe Thr AIa Val Leu Phe Ala Ala Ser Ser Aia Leu Aia

GCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CCA GGT CAG TTG TGA TGC TGT TCA CTT GCA CCG GTT TAA GGC CGA CTT CCA CAG \$\lambda

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CAA CAA AAC GGT AAA AGG TTG blie Gly Tyr Leu Asp Leu Glu Gly Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn

AGC ACA AAT AAC GGG TTA TTG TTT ATA AAT ACT ACT ATT GCC AGC ATT GCT GCT AAA TCG TGT TTA TTG CCC AAT AAC AAA TAT TTA TGA TCA TAA CGG TCG TAA CGA CCA TTT Ser Tr Asn Asn Gly Leu Leu Phe 11e Asn Thr Thr 11e A1a Ser 11e A1a A1a Lys

(hal KPI(-4-57; I16H, S17F)

Rsrll

BamHI HindIII

ACT GAA GAG TAC TOC ANG GCA GYC TOC GGA TYC GCT ATT TAA GCT T TGA CTT CYC ANG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A FTR Glu Glu Tyr Cys Met Ala Vai Cys Gly Ser Ala Ile

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 16

pTW6184

α-factor

ATG AGA TIT CCT TCA ATT TIT ACT GCA GIT TTA TIC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA Met Arg Phe Pro Ser IIe Phe Thr Ala Val Leu Phe Ala Ala Ser Ser Ala Leu Ala

ATC GOT TAC TTA GAT TTA GAA GGG GAT TTC GAT GIT GCT GIT TTG CCA TIT TCC AAC TAG CCA ATG AAT CTA AAT CIT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTG ▶ lie Gly Tyr Leu Asp Leu Glu Gly Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn

AGC AGA AAT AAC GGG TTA TIG TTT ATA AAT ACT ACT ATT GGC AGC ATT GCT GCT AAA TGG TGT TTA TTG CCC AAT AAC AAA TAT TTA TGA TGA TGA TGG TGA TGG GA GGA TTT > Ser Thr Asn Asn Gly Leu Leu Phe II e Asn Thr Thr II e Ala Sa II e Ala Sar II e Ala Ala Lys

KPI(-4-57; I16H, S17Y) Yhai

Rsrli Agel

Aatii

GCT GAG ACC GGT CCG TGC CGT GCA ATG CAC TAC CGC TGG TAC TTT GAC GTC ACT GAA CGA CTC TGG CCA GGC ACG GCA CGT TAC GTG ATG GCG ACC ATG AAA CTG CAG TGA CTT PAIA GIU The GIV Pro Cvs And Ala Met His Tve And Trp Tve Phe Asp Val The Giu

GGT ANG TGC GGT CCA TTC TTT TAC GGC GGT TGC GGC GGC AAC CGT AAC AAC TTT GAC CCA TTC ACG CGA GGT AAG AAA ATG CCG CCA ACG CCG TTG GCA TTG TTG AAA CTG FGIY Lys Cys Ala Pro Phe Phe Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp BamHi

ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A Thr Gu Giu Tyr Cys Met Ala Val Cys Giy Ser Ala Ile

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 17

pTW6185

α-	rа	C.	ſΟ	r

ATG AGA TIT CCT TCA ATT TIT ACT GCA GIT TTA TIC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA ▶Met Arg Phe Pro Ser lie Phe Thr Ala Val Leu Phe Ala Ala Ser Ser Ala Leu Ala

GCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CGA GGT CAG TTG TGA TGT TGT CTT CTA CTT TGC CGT GTT TAA GGC CGA CTT CGA CAG Ala Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gin Lie Pro Ala Glu Ala Val

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTG
▶iie Giy Tyr Leu Asp Leu Giu Giy Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn

acc aca aat aac gog tha tig tit ata aat act act att goc acc att got got aaa tog tut tha tig coc aat aac aaa tan tib tig tib ata cog too taa coa coa tit best hat as as got leu leu phe he he as int the til see the Ala ala Lys

KPI(-4-57; I16H, S17W)

GAA GAA GGG GTA TOT CTA GAT AAA AGA GAG GTT GTT AGA GAG GTG TGC TCT GAA CAA CTC CTC CCC CAT AGA GAT CTA TTT TCT GC CAA CAA TCT CTC CAC AGA AGA GCA CTC GTT GGU GGU GGU GGU VAI SAF LGU ASA LYS ATG GGU VAI ATG GGU VAI Cys SAF GGU GTA

Rsrii

Agel

GGT AAG TGC GCT CCA TTC TTT TAC GGC GGT TGC GGC GGC AAC CGT AAC AAC TTT GAC CCA TTC ACG CGA GGT AAG AAA ATG CCG CCA ACG CCG CTG GCA TTG TTG AAA CTG FGIY Lys Cys Ala Pro Phe Phe Tyr Gly Gly Cys Gly Gly Asn Arg Asn Asn Phe Asp BamHI

ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A Thr Glu Glu Tvr Cvs Met Ala Val Cvs Glv Ser Ala ile

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 18

pTW6173

α-factor

ATG AGA TIT CCT TCA ATT TIT ACT GCA GTT TTA TIC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT CGT AGG AGG CGT AAT CGA Met Arg Phe Pro Ser Ile Phe Thr Ala Val Leu Phe Ala Ala Ser Ser Ala Leu Ala

GCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CGA GGT CAG TTG TGA TGT TGT CTT CTA CTT TGC CGT GTT TAA GGC CGA CTT CGA CAG Ala Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gin II e Pro Ala Glu Ala Val

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GIT GCT GTT TTG CCA TTT TCC AAC TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTG File Gly Tyr Leu Asp Leu Glu Gly Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn

KPI(-4-57: M15A, I16H)

Rsrll

Agel Aatii GOT GAG ACC GGT CCG TGC CGT GCA GCT CAC TCC CGC TGG TAC TTT GAC GTC ACT GAA CGA CTC TGG CCA GGC ACG CCA CGT CGA GTG AGG GGG ACC ATG AAA CTG CAG TGA CTT AAI GIU Thr GIy Pro Cys Arg Ala Ala His Ser Arg Trp Tyr Phe Asp Val Thr Giu

GGT AMG TGC GCT CCA TTC TTT TAC GGC GGT TGC GGC GGC AAC GGT AAC TTT GAC CCA TTC AGC GGA GGT AAA AAA ATG GCG CCA AGG GGG CGG CCG TTG GCA TTG TAT TTG AAA CTG $\$ Giy Lys $\$ Gy $\$ Aia $\$ Pro $\$ Phe $\$ Phe $\$ Tyr Giy Giy $\$ Giy Giy Asa $\$ Arg $\$ Asa Arg $\$ Asa Asa $\$ Ara Asa $\$ Asa $\$

BamHI HindIII

ACT GAA GAG TAC TOC ATG GCA GTG TGC GGT ACC GCT ATT TAA GCT TAGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA ATT CGA ATT TAA ACT TGA CTT CTC ATG ACG TAC ACG CGT AGG GCA AAA ATT CGA A

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 19

pTW6174

α-factor

ATG AGA TIT COT TOA ATT TIT ACT GOA GIT TIA TIC GOA GOA TOC TOC GOA TIA GOT TAC TOT AAA GGA AGI TAA AAA TGA CGI CAA AAI AAG CGI COT AGG AGG CGI AAI CGA Met Arg Phe Pro Ser IIe Phe Thr Ala Val Leu Phe Ala Ala Ser Ser Ala Leu Ala

GCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CGA GGT CAG TTG TGA TGT TGT CTT CTA CTT TGC CGT GTT TAA GGC CGA CTT CGA CAG Ala Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gin Lie Pro Ala Glu Ala Val

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC TAG CCA ATG AAT CTA AAT CTT CCC CTA AAG CTA CAA CGA CAA AAC GGT AAA AGG TTG File Giv Tvr Leu Asp Leu Giu Giv Asp Phe Asp Val Ala Val Leu Pro Phe Ser Asn

KPI(-4-57: M15L, I16H)

Rsrii

Agel

OCT GAG ACC GGT CCG TGC CGT GCA TTG CAC TCC CGC TGG TAC TTT GAC GTC ACT GAA
CGA CTC TGG CCA GGC ACG GCA CGT AAC GTG AGG GCG ACC ATG AAA CTG CAG TGA CTT Ala Giu Thr Gly Pro Cys Arg Ala Leu His Ser Arg Trp Tyr Phe Asp Val Thr Glu

GGT AAG TGC GCT CCA TTC TTT TAC GGC GGT TGC GGC GGC AAC CGT AAC AAC TTT GAC CCA TTC ACG CGA GGT AAG AAA ATG CCG CCA ACG CCG CCG TTG GCA TTG TTG AAA CTG FGIY Lys Cys Ala Pro Phe Phe Tyr Giy Giy Cys Giy Giy Asn Arg Asn Asn Phe Asp BamHI

ACT GAA GAG TAC TGC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC ACG CCT AGG CGA TAA ATT CGA A

Thr Glu Glu Tyr Cys Met Ala Val Cys Gly Ser Ala ile

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 21

KPI(-4-57; M15A, S17Y) TW6166

Glu - Val - Val - Arg - Glu - Val - Cys - Ser - Glu - Gln - Ala -

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Title: PROTEASE INHIBITOR PEPTIDES Inventor(s): R. Tyler WHITE et al.

Appl. No.: 09/201,715

Figure 24

KPI(-4-57; I16H, S17F) TW6183

Figure 25

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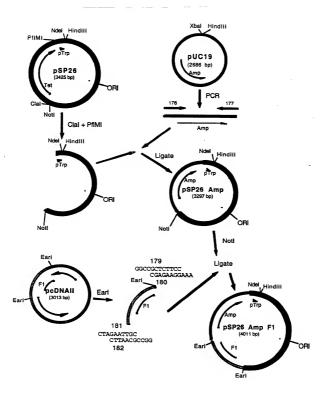
Title: PROTEASE INHIBITOR PEPTIDES Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 26

Inventor(s): R. Tyler WHITE et al.

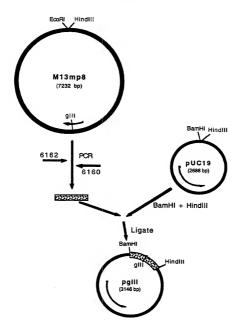
Figure 27 Appl. No.: 09/201,715

Figure 30



Title: PROTEASE INHIBITOR PEPTIDES
Inventor(s): R. Tyler WHITE et al.
Appl. No.: 09/201,715

Figure 31



Title: PROTEASE INHIBITOR PEPTIDES
Inventor(s): R. Tyler WHITE et al.
Appl. No.: 09/201,715

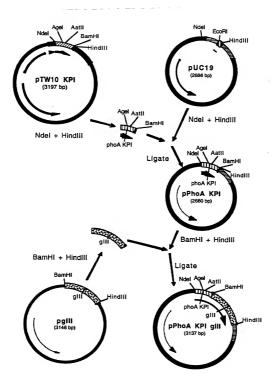
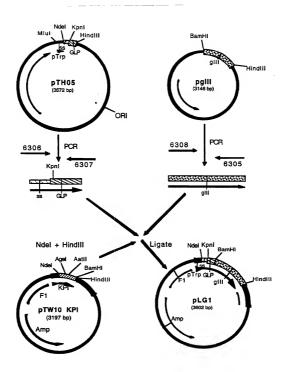
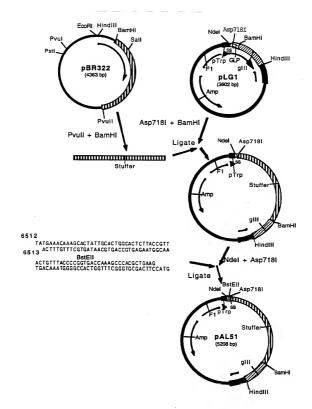


Figure 33



Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 34



Title: PROTEASE INHIBITOR
PEPTIDES
Inventor(s): R. Tyler WHITE et al.
Appl. No.: 09/201,715

Figure 35

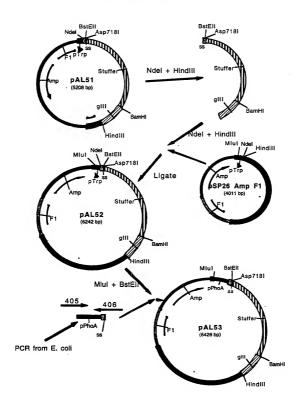


Figure 36

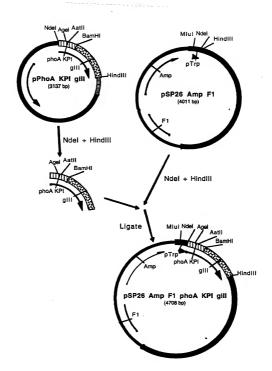
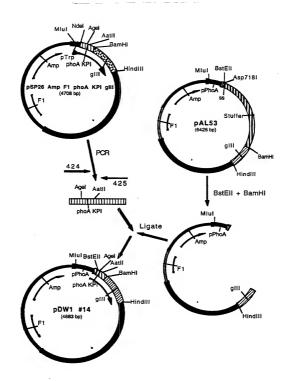


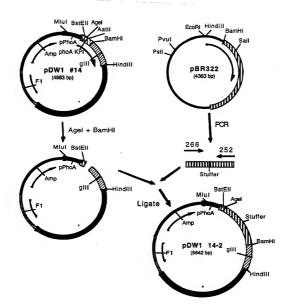
Figure 37 Appl. No.: 09/201,7

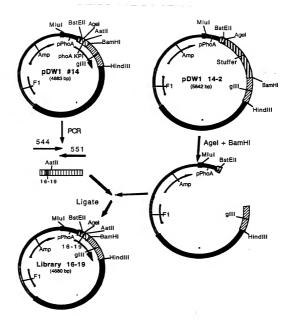


Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

phoA signal	_		B.451
GTG AAA CAA AGC	ACT ATT GCA CTG O	CA CTC TTA CCG	BstEil TTA CTG TTT ACC CCG GTG ACC AAA
Val Lys Gin Ser	Thr IIe Ala Leu A	la Leu Leu Pro	Leu Leu Phe Thr Pro Val Thr Lys
KPI (1-55)			
		Agel	
Ala Glu Val Cvs	Ser Glu Gin Ala G	HU The Gly Pro	TGC CGT GCA ATG ATC TCC CGC TGG Cys Arg Ala Met IIe Ser Arg Trp
-		,	oyo mig mice into our mig tip
Aatii			
TAC TIT GAC GIC :	ACT GAA GGT AAG 1	TOC GCT CCA TTC	TTT TAC GOC GOT TGC GOC GOC AAC Phe Tyr Gly Gly Cys Gly Gly Asn
Tyr the Asp var	ini die diy Lys c	ys Ala Più Pile	
			BamHI gili
CGT AAC AAC TIT	GAC ACT GAA GAG T	TAC TOC ATG GCA	GIG TGC GGA TCCTGGT GGT GGC TCT
PARG ASH ASH PHE	Asp inr Giu Giu i	yr Cys Met Ala	Val Cys Gly Ser [®] Gly Gly Gly Ser
			GCT AAT AAG GGG GCT ATG ACC GAA
Gly Ser Gly Asp	Phe Asp Tyr Glu L	ys Met Ala Asn .	Ala Asn Lys Gly Ala Met Thr Glu
AAT GCC GAT GAA .	AAC GCG CTA CAG T	CT GAC GCT AAA	GGC AAA CIT GAT TOT GTC GCT ACT
▶Asn Ala Asp Glu	Asn Ala Leu Gin S	Ser Asp Ala Lys	Gly Lys Leu Asp Ser Val Ala Thr
GAT TAC GGT GCT	GCT ATC GAT GGT 1	TTC ATT GGT GAC	GTT TCC GGC CTT GCT AAT GGT AAT
			Val Ser Gly Leu Ala Asn Gly Asn
com com som com	01 m mm com coo n		ATG GCT CAA GTC GGT GAC GGT GAT
			Met Ala Gin Val Giv Asp Giv Asp
			CCT TCC CTC CCT CAA TCG GTT GAA Pro Ser Leu Pro Gin Ser Val Glu
TGT CGC CCT TTT	GTC TTT GGC GCT G	GT AAA CCA TAC	GAA TIT TOT ATT GAT TGT GAC AAA
FCys Arg Pro Phe	vai Phe Gily Ala G	∄ly Lys Pro Tyr	Glu Phe Ser IIe Asp Cys Asp Lys
ATA AAC TTA TTC	CGT GGT GTC TTT C	CG TTT CTT TTA	TAT GIT GCC ACC TIT ATG TAT GTA
▶ lle Asn Leu Phe	Arg Gly Val Phe A	la Phe Leu Leu	Tyr Val Ala Thr Phe Met Tyr Val
TTT TCT ACG TTT	OCT AAC ATA CTG	TOT A AT AAG CAC	ጥጥ ጥልል ጥል
Phe Ser Thr Phe			

Figure 39





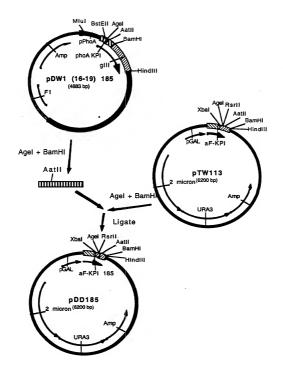
Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

phoA signal			
	>		BstEII
			CTG TTT ACC CCG GTG ACC AAA
▶Val Lys Gin Ser Th	r lie Ala Leu Ala	Leu Leu Pro Leu	Leu Phe Thr Pro Val Thr Lys
KPI (1-55; 1	6 - 19)	Agel	16 - 19
		Agei Lace centree mee	CGT NNS NNS NNS TGG TAC
			Arg Trp Tyr
		,, .	
Aatii			
TTT GAC GTC ACT GA	A GGT AAG TGC GCT	CCA TTC TTT TAC	GGC GGT TGC GGC GGC AAC CGT
▶ Phe Asp Val Thr GI	u Gly Lys Cys Ala	Pro Phe Phe Tyr	Gly Gly Cys Gly Gly Asn Arg
			BamHi Gili
			GGA TCC GGT GGT GGC TCT GGT
PASH ASH PHE ASP II	r Giu Giu iyr Cys	Met Ala val Cys	Gly Ser Gly Gly Ser Gly
THE COT CAST TITTE CO	ידים מבה בהם ידים ידי	CCA AAC CCT AAT	AAG GGG GCT ATG ACC GAA AAT
			Lvs Giv Ala Met Thr Giu Asn
,,			-,,
			CTT GAT TCT GTC GCT ACT GAT
≯Ala Asp Glu Asn Al	a Leu Gin Ser Asp	Ala Lys Gly Lys	Leu Asp Ser Val Ala Thr Asp
mag com com com a	va cam com mma am		GGC CTT GCT AAT GGT AAT GGT
			Gly Leu Ala Asn Gly Asn Gly
· iyi Giy Ala Ala ii	o Asp City The Tit	, di , Aup 141 001	מוץ בספ אום אטוו מוץ אטוו מוץ
			gIII
GCT ACT GGT GAT T	T GOT GGC TOT AA	TCC CAA ATG GCT	CAA GTC GGT GAC GGT GAT AAT
▶Ala Thr Gly Asp Pt	e Ala Giy Ser Ası	Ser Gin Met Ala	Gin Val Giy Asp Giy Asp Asn
mas com mms sma s	.mm mma .com .c.:		CTC CCT CAA TCG GTT GAA TGT
			Leu Pro Gin Ser Val Glu Cvs
Poer Pro Lea Met As	in Ash Fhe Arg Gil	i tyr Leu Pro Ser	Led Pro Gill Ser val Gid Cys
CGC CCT TTT GTC T	T GGC GCT GGT AA	CCA TAC GAA TTT	TCT ATT GAT TGT GAC AAA ATA
Arg Pro Phe Val Pt	e Gly Ala Gly Lys	Pro Tyr Glu Phe	Ser Ile Asp Cys Asp Lys Ile
-			
			GCC ACC TIT ATG TAT GTA TIT
▶Asn Leu Phe Arg G	y Val Phe Ala Phe	Leu Leu Tyr Val	Ala Thr Phe Met Tyr Val Phe
TOT ACG TIT GOT A	AC ATA CTG CGT AA	P ANG GAG TOT TAA	TA
Ser Thr Phe Ala As			
Joe IIII FIIO ATA AS	III IIO LOU AIG ASI	Lys Giu Ser 111	

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

ph	oA s	igna	t		_	-		-	-		-	-	-						
_					-												tEll		
	AAA																		
▶ Va I								Ai a	Leu	Leu	PF 0	Leu	Leu	Pne	inr	PF 0	vai	inr	Lys
	KP	1 (3-	-55;	M15	οA,	51/	-)	_	Agel										
	GAG							GAG	ACC										
▶Ala	Gl u	Val	Cys	Ser	Glu	GIn	Ala	Gш	Thr	GГу	Pro	Cys	A rg	Ala	Ala	lie	Phe	Arg	Trp
The C	TTT	Aatl		3.00	CNA	com	220	ma		CC3	mm~		mac	~~~	~~	m	ccc	ccc	330
▶ Tyr																			
,.						,	-,-	-, -					٠,,					,	
														Bam		9	111	-	
CCI	AAC	AAC	TTT	GAC	ACT	GAA	GAG	TAC	TGC	ATG	GCA	GIG	TGC	GGA	TCC	CCI	GGT	GGC	TCT
▶Arg	Asn	Asn	Phe	Asp	inr	Glu	GIU	iyr	Cys	Met	AI a	vai	Cys	GI y	Ser	GI y	GI Y	GI Y	Ser
	TCC																		
▶ GIy	Ser	GI y	Asp	Phe	Asp	Tyr	GI u	Lys	Met	Al a	Asn	Ai a	Asn	Lys	GI y	Ala	Met	Thr	Glu
3.50	GCC	Cam	CAA	220	~~	~ma	CNC	TV~TI	arc	COTT	222	ccc	***	CHAIL.	CAT	TV-VIII	carc.	COL	a.cer
P Asn																			
													•						
	TAC																		
▶Asp	lyr	GI y	Ala	Ala	110	ASP	Giy	Phe	110	GIY	Asp	vai	Ser	Gi y	Leu	Ala	ASI	Giy	ASI
GGT	GCT	ACT	GGT	GAT	TTT	GCT	GGC	TCT	AAT	TCC	CAA	ATG	GCT	CAA	GTC	GGT	GAC	GGT	GAT
▶ GI y	Ala	Thr	GI y	Asp	Phe	Ala	GI y	Ser	Asn	Ser	Gln	Met	AI a	GIп	Val	GI y	Asp	GI y	Asp
	TCA		mma.	3000		330	mma	~~	C2.2	mam	mma		m	~~~	com	CNA	·	com.	Cha
AAI																			
										•									
	CGC																		
▶ Cys	Arg	Pro	Phe	Val	Phe	GI y	Ala	GГу	Lys	Pro	Tyr	Glu	Phe	Ser	He	Asp	Cys	Asp	Lys
АТА	AAC	TTA	TTC	CGT	GGT	GTC	TTT	GCG	TTT	CTT	TTA	TAT	GTT	GCC	ACC	TTT	ATG	TAT	GTA
▶11e																			
~~~	TCT	300				ama	~~~	~~	330		C3.C	man.	ma a	mx					
▶ Phe													LAA	LA					
. / 110	-0.							9		-,,		-0.							

Figure 43



Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201.715

Figure 44

pDD185

~-1	2	ct	^	,

ATG AGA TIT CCT TCA ATT TIT ACT GCA GTT TTA TTC GCA GCA TCC TCC GCA TTA GCT TAC TCT AAA GGA AGT TAA AAA TGA CGT CAA AAT AAG CGT GGT AGG AGG GGT AAT CGA PMet Arg Phe Pro Ser IIe Phe Thr Ala Val Leu Phe Ala Ala Ser Ser Ala Leu Ala

GCT CCA GTC AAC ACT ACA ACA GAA GAT GAA ACG GCA CAA ATT CCG GCT GAA GCT GTC CCA GGT CAG TTG TCA TCAT TCT CTT CTC CCT GTT TTA GGC CGA CTT CGA CAG AIA Pro Val Asn Thr Thr Thr Glu Asp Glu Thr Ala Gln lie Pro Ala Glu Ala Val

ATC GGT TAC TTA GAT TTA GAA GGG GAT TTC GAT GTT GCT GTT TTG CCA TTT TCC AAC TAG CCA ATA TCTA AAT CTT ACT CTC CTTA AAG CTTA AAC GAA AAA GAG GTTA BHILE GIJ Tyr Leu Asp Leu Giu Giy Asp Phe Asp Vai Aia Vai Leu Pro Phe Ser Asn

AGC ACA ANT AAC GGG TTA TIG TIT ATA ANT ACT ACT ATT GGC AGC ATT GCT GGT AAA TGG TGT TTA TIG CCC AAT AAC AAA TAT TTA TGA TGA TAA GGG TGT AC GG GGA TIT > Ser Thr Asn Asn Gy Leu Leu Phe lie Asn Thr Thr Ile Ala Ser Ile Ala Ala Lys

KPI(-4-57; M15A, S17F)

#### Xhal

GAA GAA GGG GTA TOT CTA GAT AAA AGA GAG GTT GTT AGA GAG GTG TOC TOT GAA CAA CTT CTT CCC CAT AGA GAT AGA TTT TCT CTC CAA CAA ATCT CTC CAA CAG AGA CTT GTT >Gu Glu Gly Val Ser Leu Asp Lys Arg Gu Val Val Arg Glu Val Val Ser Glu Glin

#### Rsrii

#### Agel

#### BamHI Hindli

ACT GAA GAG TAC TOC ATG GCA GTG TGC GGA TCC GCT ATT TAA GCT T TGA CTT CTC ATG ACG TAC CGT CAC ACG CCC TAGG CGA TAA ATT CGA A > Thr Glu Gu Tyr Cys Met Ala Vai Cys Gly Ser Ala Ile Figure 45

Title: PROTEASE INHIBITOR PEPTIDES
Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

#### Plasma kallikrein inhibition by KPI (-4-57) variants

Varian	<u>t</u>		Substitution		K(nM)
		15	16	17	
TW11:	3 KPI (-4-57)				45.00
DD185	5 KPI (-4-57; M15A, S17F)	A		F	0.39
TW61	65 KPI (-4-57; M15A, S17W)	Α		W	0.65
TW61	66 KPI (-4-57; M15A, S17Y)	Α		Y	0.40
TW61	75 KPI (-4-57; M15L, S17F)	L		F	0.50
BG02	8 KPI (-4-57; M15L, S17Y)	L		Y	1.10
TW61	83 KPI (-4-57; I16H, S17F)		H	F	1.20
TW61	84 KPI (-4-57; I16H, S17Y)		H	Y	0.91
TW61	85 KPI (-4-57; I16H, S17W)		H	W	1.30
TW61	73 KPI (-4-57; M15A, I16H)	Α	H		1.00
TW61	74 KPI (-4-57; M15L, I16H)	L	H		0.90

			Inhibition Ki (nM)	Ki (nM)	
Variant	Sequence	kallikrei	Plasmin	XIIa	Xa
Aprotinin	RPDFCLEPPYTGPCKARI I RYFYNAKAGLCQTFVYGGCRAKRNNFKSAEDCMRTCGGA	20.00	0.23	2000.0	
Aprotinin R15, S42	DPCLEPPYTGPCRARIIRY PYNAKAGLOQTFVYGGCRAKSNNFKSAEDCMRTCGGA	0.91	0.17	3983.0	
KPI (4-57)	EVVREVCSEQAETGPCRAMISRWYFDVTEGRCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	42.00	34.00	3718.0	161.0
TW6167	EVVREVCSEQAEPGPCRAMISRWY FDVTEGRCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	61.00		3641.0	288.0
BG031	EVVREVCSEQAEVGPCRAMI SRMY FDVTEGRCAPFFYGGCGGNRNN FDTEEY CMAVCGSAI	34.00		498.0	
BG032	EVVREVCSEQAESGPCRAMISRMY FDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	49.00		731.0	
TW101	EVCSEQAETGPCKAMI SRWY PDVTEGKCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	2000.00	11.50		
TW6208	EVVREVCSEQAETGPCRGMI SRWY FDVTEGKCAPFFYGGCGGNRNN FDTEEY CMAVCGSAI			369.0	
TW106	EVCSEQAETGPCRARI SRWY FDVTEGKCAPFFYGGCGGNRNN FDTEEY CMAVCGSAI	\$60.00	3.70		
DD108	EVVREVCSEQAETGPCRAAISRWY PDVTEGKCAPFYGGCGGNRNN PDTEEY CMAVCGSAI	1.70	11.20	1600.0	123.0
DD109	EVVREVCSEQAETGPCRAIISRMY PDVTEGKCAPFFY GGCGGNRNN PDTEEY CMAVCGSAI	9.50		1681.0	421.0
DD110	EVVREVCSEQAETGPCRALISRMY FDVTEGRCAPFFY GGCGGNRNNFDTEEY CMAVCGSAL	2.10		624.0	55.0
DDIII	BVVREVCSEQAETGPCRASI SRWY FDVTEGKCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	2.60			
DD112	EVVREVCSEQAETGPCRAVISRMY PDVTEGRCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	6.80		998.0	
TW6179	EVVREVCSEQAETGP CRAGISRMY PDVTEGK CAPPTYGG CGGNRNN PDTEEY CMAVCGSAI	78.00		368.0	
TW6163	EVVREVCSEQAETGPCRAMHSRMY PDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	4.70	103.58	4532.0	457.0
TW6172	EVVREVCSEQAETGPCRAMASRWY PDVTEGRCAPFFY GGCGGNRNN FDTEEY CMAVCGSAI	315.00			1463.0
TW6180	EVVREVCSEQAETGPCRAMFSRWY FDVTEGKCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	70.00		885.0	39.0
TW6181	BVVREVCSEQAETGPCRAMKSRWYFDVTEGRCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	150.00		1514.0	
BG001	EVVREVCSEQAETGPCRAMLSRWYFDVTEGRCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	38,00	10.00	489.0	204.0
TW116	EVCSEQAETGPCRAMI IRWY FDVTEGRCAPFFYGGCGGNRHNFDTEEY CMAVCGSAI	145.00	89.00		806.0
DD 102	EVVREVCSEQAETGPCRAMIPRMY PDVTEGKCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	16.00		315.0	
DD 103	EVVREVCSEQAETGPCRAMI PRMY PDVTEGKCAP F F Y GGCGGNRHN PDTEBY CMAVCGSAI	17.00		2128.0	110.0
DD104	EVVREVCSEQAETGPCRAMIYRWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	15.00		237.0	345.0
DD105	<b>EVVREVCSEQAETGPCRAMIWRWYFDVTEGRCAPPPYGGCGGNRNNFDTEEYCMAVCGSAI</b>	18.00		198.0	320.0
TW6168	EVVREVCSEQAETGPCRAMILRWY FDVTEGRCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	25.80		3521.0	395.0
TW6182	EVVREVCSEQAETGPCRAMIHRWY FDVTEGRCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	36.00		752.0	
TW6194	EVVREVCSEQAETGPCRAMI ERMY PDVTEGRCAPFFYGGCGGNRNNFDTEEY CMAVCGSAI	70.83			
TW6210	EVVREVCSEQAETGPCRAMIQRHY PDVTEGKCAPFFYGGCGGNRHNFDTEEY CMAVCGSAI	54.00		277.0	
CL006	EVVREVCSEQAETGPCRAMI SAWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	110.20		89600.0	133.0
BG012	EVVREVCSEQAETGPCRAMISTWY FDVTEGRCAPFFY GGCGGNRNNFDTEEY CMAVCGSAI			40.0	116.0

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Title: PROTEASE INCLUDIOS PEPTIDES Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

## FIGURE 46B

			l		
TW6209	EVVREVCSEOAETGPCRAMISHWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	81.00	45.90	184.0	613.0
TW6311	EVUREVCSEOARTGPCRAMISKWY FDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	184.00		402.0	
DD128	EVVRRVCSEOAETGPCRAMISLWYFDVTBGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	44.00			37.0
TW6142	EVVRRVCSEOAETGPCRAMISRMYFDVTEGKCAPFVYGGCGGNRNNFDTEEYCMAVCGSAI	18.00	18.00	7972.0	225.0
AI 301	EVVRRUCSROAFTGPCRAMISBWYFDVTEGKCAPFLYGGCGGNRNNFDTEEYCMAVCGSAI	216.00		1557.0	
AI 302	RIVERVICERDARTGPCRAMISRAYEDVTEGKCAPFGYGGCGGNRNNFDTEEYCMAVCGSAI	39.00			316.0
TW6147	RUVERUCSEDARTOPCRAMISRWYPDVTEGKCAPFFYGGCAGNRNNFDTEEYCMAVOGSAI	35.00		1090.0	179.0
TW6138	RUMBRUCSROAFTGPCRAMISRWYPDVTEGKCAPFFYGGCKGNRNNFDTEEYCMAVCGSAI	18.00		921.0	309.0
TW6154	RUVERUCSEDAETGPCRAMISRWYPDVTEGKCAPFFYGGCLGNRNNFDTEEYCMAVCGSAI	11.00		915.0	39.0
TW6155	RUVERUCSEOAETGPCRAMISRMYPDVTEGKCAPFFYGGCMGNRNNFDTEEYCMAVCGSAI	11.00			27.0
TW6140	EVVREVCSEOAETGPCRAMISRWYFDVTEGKCAPFFYGGCNGNRNNFDTERYCMAVCGSAI	35.00		475.0	
TW6156	EVVREVCSEORETGPCRAMISRMY PDVTEGRCAPFFYGGCPGNRNNFDTEEYCMAVCGSAI				
TW6141	RUVRRUCSEOAETGPCRAMISRWYPDVTEGRCAPFFYGGCQGNRNNFDTEEYCMAVCGSAI	42.00			
TW118	EVCSEOAETGPCRAMISRWYFDVTEGKCAPFFYGGCRGNRNNFDTEEYCMAVCGSAI	9009	24.00	13009.0	089
00100	EVVREVCSEOAETGPCRAMISRWY FDVTEGKCAP FFYGGCCGNRNN FDTEEYCMAVCGSAI	15.00			
TW6157	EVVREVCSEORETGPCRAMISTRY FDVTEGKCAPFFYGGCSGNRNNFDTEEYCMAVGSAI	40.00		511.0	168.0
TW6158	EVVREVCSEOAETGPCRAMISRWYPDVTEGRCAPFFYGGCTGNRNNPDTERYCMAVCGSAI	29.00			
TW6159	EVVREVCSEOAETGPCRAMISRWY PDVTEGRCAPFFY GGCVGNRNN PDTEEY CMAVCGSAI	17.00			64.0
TWKIKI	EVVREVCSEOAETGPCRAMISRWYPDVTEGKCAPFFYGGCYGNRNNFDTEEYCMAVCGSAI	7.50	18.00	1507.0	8.7
וסוממ	EVVREVCSEOAETGPCRAMISRWYFDVTEGKCAPFYGGCDGNRNNFDTEEYCMAVCGSAI	64.00		924.0	
TW6151	RVVREVCSEORETGPCRAMISRMYFDVTEGKCAPFFYGGCEGNRNNFDTEEYCMAVCGSAI	163.00		1162.0	954.0
TW6139	EVVREVCSEORETGPCRAMISRNY FDVTEGKCAPFFYGGCHGNRNNFDTEEYCMAVCGSAI	19.00	22.80	152.0	78.0
TW6153	EVVREVCSEORETGPCRAMISRWYPDVTEGKCAPPFYGGCIGNRNNFDTEEYCMAVCGSAI	11.20	21.30	65.0	36.0
TW122	EVCSEQAETGPCRAMISRWYFDVTEGKCAPFFYGGCGANRNNFDTEEYCMAVCGSAI	32.00	27.00		581.0
TW6178	BUVREVCSEQAETGPCRAMISRWYFDVTEGRCAPFFYGGCGRNRNNFDTEEYCMAVCGSAI	16.00		444.0	
TW6148	EVVREVCSEQAETGPCRAMI SRWY PDVTEGKCAPPY YGGCGGARNN PDTEEY CMAVCGSAI	40.00	_		
TW124	EVCSEQAETGPCRAMISRWYPDVTEGRCAPFFYGGCGGNSNNFDTEEYCMAVCGSAI	64.00	48.00		
TW6149	EVVREVCSEQAETGPCRAMISRWY PDVTEGKCAPFFYGGCGGNANNFDTEEYCMAVCGSAI	24.00			
TW6173	EVVREVCSEQAETGPCRAAHSRWY PDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	8.		1432.0	
TW6174	EVVREVCSEQAETGPCRALHSRWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	8.0		2796.0	
BG002	EVVREVCSEQAETGPCRALLSRWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	0.98	19.00	403.0	00.0
DD129	EVVREVCSEQAETGPCRALFSRWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	3.60		1864.0	0.0
DD185	EVVREVCSEQAETGPCRAAIFRWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	0.39	8.71	150.0	196.0

Tride: PROTEASE INHIBIT ON PEPTIDES

PEPTIDES

PEPTIDES

APPL No: 09/201,715

## FIGURE 46C

EVIRENCESEQUETORIAL TIANY POLTEGICA PETGOCOGININI POTESE CIAN COSAL  EVIRENCESEQUETORIAL LIANY POLTEGICA PETGOCOGININI POTESE CIAN COSAL  EVIRENCESE ATORIA POR LIANY POLTEGICA PETGOCOGININI POTESE CIAN COSAL  EVIRENCESEQUETORIAL LIANY POLTEGICA PETGOCOGININI POTESE CIAN COSAL  EVIRENCESEQUETORIAL LIANY POLTEGICA PETGOCOGININI POTESE CIAN COSAL  EVIRENCESEQUETORIAL LIANY POLTEGICA PETGOCOGININI POTESE CIAN CORANI PORTEGICA CIAN COSAL  EVIRENCESEQUETORIAL LIANY POLTEGICA PETGOCOGINI PORTEGICA CIAN CORANI P				L	I	
EVIRENCESQUETGERALI INTRIPRIVEDATEGROAPE PROGOGGHINIPEDTERICANUCGSAI  EVIRENCESQUETGERALI INTRIPRIVEDATEGROAPE PROGOGGGHINIPEDTERICANUCGSAI  EVIRENCESQUETGERALI INTRIPRITEGROAPE PROGOGGGHINIPE	TW6165	EVVREVCSEOAETGPCRAAIWRMY PDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	0.65		206.0	
EVAREACESQUETO PERALLITRINE POVEGRCARP PF VGG COGNINNE PER CHANCGSAIL  EVAREACESQUETO PERALLITRINE POVEGRCARP PF VGG COGNINNE PER	TW/6166	PULD PULCEDAR PROPERTY PROPERTY CAPPET GCCGCONNINFOTERY CMANCGSAI	0.4	10.10	73.0	
EVAREACSEQUETO PERAL LI GANT PROVEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN PDTEET CHANCOS A.I.  EVAREACSEQUETO PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOG PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOG PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOG PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOG PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOG PERAL LI GANT POVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOGGORAL LI TRAY TOVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVAREACSEGAL TOGGORAL LI TRAY TOVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVARBUCESEGAL TOGGORAL LI TRAY TOVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVARBUCESEGAL TOGGORAL LI TRAY TOVTEGKCAR PF TOGGOGGINAIN POTE EXCHANCOS A.I.  EVARBUCESEGAL TOGGORAL POTEGRA TOR PT TOGGOGGINAIN POTEGRA TORANCOS A.I.  EVARBUCESEGAL TOGGORAL TRAY TOVTEGKCAR PF TOGGOGGINAIN POTEGRA TORANCOS A.I.  EVARBUCESEGAL TOGGORAL TRAY TO	1 10000	PHINDENCEDAR TO THE WAY PROTECT CAPPEY GCCCONRUN POTEET CHAVCGSAI	1.10	12.10	93.8	
EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PTEET CHANCOS AL  EVAREACES QUESTO PERALLI FRAY POYTEGACARP PT GGGGGGRINN PT	BC028	EVYREVCESQUELGE CRAIL LINE TO THE CONTROL OF THE CO	1.20		0.619	111.0
EVAREACESQUETO PERMATERIA POVTEGRCAP PF TOGGOGNINNEDTER TCHANCGSAI  EVAREA	1 W0107	#10TO BELLEVILLE CONTROL TO PROVE PROVER CRAPETY GGCGGNRNNFDTEEY CMAVCGSAI	0.85	12.80	293.0	74.0
EVIRENCESQAETGCRAD I HRWIT DDYEGRCAP PYGGCGGRINNEPDTEEKCHANCGSAI  EVURENCESQAETGCRAD I HRWIT DDYEGRCAP PYGGCGGRINNEPDTEEKCHANCGSAI  EVURENCESQAETGCRAD I FRRWIT DDYEGRCAP PYGGCGGRINNEPDTEEKCHANCGSAI  EVURENCESQAETGCRAD I FRRWIT DDYEGRCAP PYGGCGGRINNEPDTEEKCHANCGSAI  EVURENCESQAETGCRAD I SIRWIT DDYEGRCAP PYGGCGGGRINNEPDTEEKCHANCGSAI  EVURENCESQAETGCRAD I SIRWIT PDYTEGRCAP PYGGCGGGRINNEPDTEEKCHANCGSAI  EVURENCESGAETGCRAD I SIRWIT PDYTEGRCAP PYGGCGGGRINNEPDTEEKCHANCGSAI  EVURENCESGAETGCRAD I SIRWIT PDYTEGRCAP PYGGCGGGRINNEPDTEEKCHANCGSAI  EVURENCESGAETGCRAD SIRWIT PDYTEGRCAP PYGGCGGGRINNEPDTEEKCHANCGSAI  EVURENCESGAET	TUX176	PHYMAN CORPORATION TO THE WAY POUTEGROAP PYGGOGGINNIN POTERY CANAVOGSAI	0.50	7.46	35.0	96.0
EVIRENCES PARTOCRAGE INNET DOTES CAN PER TOGGOGORNIN POTES CHANCOS A. I  EVURENCES PARTOCRAGE HENTE DOTES CAN PER TOGGOGORNIN POTES CHANCOS A. I  EVURENCES PARTOCRAGE HENTE DOTES CAN PER TOGGOGORNIN POTES CHANCOS A. I  EVURENCES PARTOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PARTOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IS MAY TO TESCALA PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IN PROTES CHAN PET POTGOGOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IN PROTES CHANCOR PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IN POTTES CHANCOR PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IN POTTES CHANCOR PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IN POTTES CHANCOR PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOCRAM. IN POTTES CHANCOR PET VOGCOGORNIN POTES CHANCOS A. I  EVURENCES PATOR POTES MAY TO TESCALA PET POTGOGOGORNIN POTES CHANCOS A. I  EVURENCES PATOR POT POT THE MENTO POT TESTA PET P	TWEAD!	WINDSHOCKPARTCHCOCK TO THE TOTAL THE	34.60		419.0	
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EVIPRECESQUETGERALISMY PDVTEGRCAP PFOGGOGNRIN PDTEET CHANGGSAI  EVAREVCESQUETGERALISMY PDVTEGRCAP PFOGGOGNRIN PDTEET CHANGGSAI  EVAREVCESQUETGERALISMY POTTEGRCAP PFOGGOGNRIN PTEET CHANGGSAI  EVAREVCESQUETGERALISMY POTTEGRCAP PFOGGOGNRIN PROTTEET CHANGGSAI  EVAREVCESQUETGERALISMY POTTEGRCAP PFOGGOGNRIN PROTTEET CHANGGSAI  EVAREVCESQUETGERALISMY POTTEGRCAP PFOGGOGNRIN PTEET CHANGGSAI  EVAREVCESQUETGERALISMY POTTEGRCAP PFOGGOGNRIN PTEET CHANGGSAI  EVAREVCESCONG PROTTEGRCAP PFOGGOGNRIN PROTTEET CHANGGSAI  EVAREVCESCONG PFOGGOGNRIN PROTTEGRCAP PFOGGOGNRIN PROTTEET CHANGGSAI  EVAREVCESCONG PFOGGOGNER PFOGGOGNRIN PROTTEET CHANGGSAI  EVAREVCESCONG PFOG	TW6203	EVVREUCERDA ETGPCRAGIPRAY FDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	31.20		5045.0	
EVIRENCESQUETO COLLIA MATED VEGICLA PETGGGGGRINNED PERCHANCOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGRINNED PERCHANCOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGRINNED PERCHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PERCHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGRINNED PERCHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PERECHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PERECHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PERECHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PERECHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESQUETO COLLIA LISHET DOTEGACA PETGGGGGGRINNED PERCHANGOSAL  EVURENCESQUETO COLLIA LISHET POTTEGACA PETGGGGGGRINNED PERCHANGOSAL  EVURENCESQUETO COLLIA LISHET POTTEGACA PETGGGGGGRINNED PERCHANGOSAL  EVURENCESQUETO COLLIA LISHET POTTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESQUETO COLLIA LISHET POTTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESQUETO COLLIA LISHET POTTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESCOLETO PETCHA PETRET POTTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESCOLETO PETCHA PETAT PETRET POTTEGACA PETGGGGGGRINNED PETER CHANGOSAL  EVURENCESCOLETO PETCHA PETAT P	TW2004	PHIRD WINDS TO THE TRANSPORTEGICA PETERGECONNINFOTEET CHANGES AT			147.0	87.0
EVITED CEREBALTO CORA A SENTE TO TEGE CAN PERGOCCIONINI POTE ER CHANCOS A I EVURENCES CHETO CORA A SENTE TO TEGE CAN PETGGO CONTINI POTE TEGE CHANCOS A I EVURENCES CHETO CRAMITE MAY TO TEGE CAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO CRAMITEMENT DOTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN I THAY POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN ET SHATE POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN ET SHATE POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN ET SHATE POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN ET SHATE POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN ET SHATE POTT ESCAN PETGGO CONTINI POTE EXCHANCOS A I EVURENCES CHETO ETCHAN	1 W0204	PHYSE POST CONTROL OF THE PROPERTY OF THE PROP			195.0	29.0
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EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGRINN PDTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTESCALAP PYGGCGGGRINN PPTEST CHANCOSAL  EVINENCESQUETO CRAMITERY POTTESCALAP PYGGCGGGRINN PPTEST CHANCOSAL  EVINENCESQUETO PROCESAL PYTER PROTEST PYTEST CHANCOSAL  EVINENCESQUETO PROCESAL PYTER PYTEST	DD114	SHAND STATES OF THE STATES OF	0.83	52.20	589.0	1396.0
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EVITATIOSEQUETO PERMAHERAT POVTEGICAR PETGGGGGRINNEPTEET CHANGGSAI  EVITATIOSEQUETO PERMAHERAT POVTEGICAR PETGGGGGRINNEPTEET CHANGGSAI  EVITATIOSEQUETO PERMAHERAT POVTEGICAR PETGGGGGRINNEPTEET CHANGGSAI  EVITATIOSEQUETO PERMAI TRAPT POVTEGICAR PETGGGGGRINNEPTEET CHANGGSAI  EVITATIOSEQUETO PERMAI TRAPT POVTEGICAR PETGGGGRINNEPTEET CHANGGSAI  EVITATIOSEQUETO PERMAI TRAPT POVTEGICAR PETGGGGGRINNEPTEET CHANGGSAI  EVITATIOSEGUETO PERMAI TRAPT POVTEGICAR PETGGGGGRINNEPTEET CHANGGGGAI  EVITATION PETGGGGGRINNEPTEET CHANGGGGGRINNEPTEET CHANGGGGGGRINNEPTEET CHANGGGGGGRINNEPTEET CHANGGGGGGRINNEPTEET CHANGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	1 W0103	STATES OF THE ST	16.0	11.96	14000.0	214.0
EVITATO-CSEQUETO CENAL LIGHT POLYZGICKARP PT FOGGOGNINN PDTEET CHANCOGAL  EVITATO-CSEQUETO CENAL LIGHT POLYZGICKAP PT FOGGOGNINN PDTEET CHANCOGAL  EVITATO-CSEQUETO CORRAN POLYZGICKAP PT FOGGOGNINN PDTEET CHANCOGAL  EVITATO-CSEQUETO POLYTIN POLYZGICKAP PT FOGGOGONINN PDTEET CHANCOGAL  EVITATO-CSEQUETO POLYTIN POLYZGICKAP PT FOGGOGONINN POLYZGICHOLOGAL  EVITATO-CSEQUETO POLYTIN POLYTIN POLYTIN POLYZGICHOLOGAL  EVITATO-CSEQUETO POLYTIN POLYT	1 WO104	STATES TO SECOND STATES OF THE	1.30	18.60	388.0	473.0
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EVVREVCERQUETO PETRINITRY PTOVTEGICARP PTGGCTGRININ PTERFORM VOGSAI  EVVREVCERQUETO PETRINITRY PTOVTEGICARP PTGGCGGRININ PTERFORM VOGSAI  EVVREVCERGE SETT OF IT THAT PTOVTEGICARP PTGGCGGRININ PTERFORM VOGSAI  EVVREVCERGE SETT OF PETRININ PTOVTEGICARP PTGGCGGRININ PTERFORM VOGSAI  EVVREVCERGE SETT OF PETRININ PTOVTEGICARP PTGGCGGRININ PTERFORM VOGSAI  EVVREVCERGAETO PETRININ PTOVTEGICARP PTGGCGGGRININ PTERFORM VOGSAI  EVVREVCERGAETO PETRININ PTOVTEGICARP PTGGCGGGRININ PTERFORM VOGSAI  EVVREVCERGAETO PETRININ PTOVTEGICARP PTGGCGGGRININ PTERFORM VOGSAI  EVVREVCERGAETO PETRINI PTOVTEGICARP PTGGCGGGRININ PTERFER CHANCGSAI  EVVREVCERGAETO PETRINI PTOVTEGICARP PTGGCGGGRININ PTERFER CHANCGSAI  EVVREVCERGAETO PETRINI PTOVTEGICARP PTGGCGGGRININ PTERFER CHANCGSAI  EVVREVCERGAETO PETRINI PTOVTEGICARP PTGGCGGGRININ PTERFORM VOGSAI  EVVREVCERGAETO PETRINI PTOVTEGICARP PTGGCGGGRININ PTERFORM VOGSAI  EVVREVCERGAETO PETRINI PTOVTEGICARP PTGGCGGGRININ PTERFER CHANCGGRININ PTERFORM VOGGGRININ PTERFORM	1 WOIO/	PHILIPPINGED BETGE OF THE PROPERTY CAPPET CONTRIBUTE CHANCES AT	4.00		419.0	24.0
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EVVENCESQUETO PURA JETHINET FOVTEGKCAP PTGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOVTEGKCAP PTGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOVTEGKCAP PTGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOVTEGKCAP PTGGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOVTEGKCAP PTGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOVTEGKCAP PT TGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOVTEGKCAP PT TGGGGGRINNED TEET CHANGGSAI  EVVENCESQUETO PURA JETHINET FOR THE TEET CHANGGSAI  EVVENCES THE TEET CHANGGS TH	1 401/0	PANDENCEDAR TOP			213.0	299.0
EVUENCERQUETGPCRAHFRHY PDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALALEHRY PDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALALEHRY PDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALETRAY PDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALETRAY FDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALETRAY FDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALETRAY FDVTBGKCARP FYGGCGGRRINR PDTEEVCHAVCGSAI  EVHENCESQUETGPCRALETRAY FDVTBGKCARP FYGGCGGRRINR PDTEETCHAVCGSAI  EVHENCESGUETGPCRALETRAY FDVTBGKCARP FYGGCGGRRINR PDTEETCHAVCGSAI  EVHENCESGUETGPCRALETRAY FDVTBGKCARP FYGGCGGRRINR PDTEETCHAVCGSAI  EVHENCESGUETGPCRALETRAY FFANT FORTER FFANT FORTER FYGGCGGRRINR PDTEETCHAVCGSAI  EVHENCESCHAFFART FFANT FORTER FFANT FORTER FYGGCGGRRINR PDTEETCHAVROGGSAI  EVHENCESCHAFFART FFANT FORTER FFANT FORTER FYGGCGGRRINR PDTEETCHAVROGGSAI  EVHENCESCHAFFART FFANT FORTER FFANT FORTER FYGGCGGRRINR PDTEETCHAVROGGAI  EVHENCESCHAFFART FFANT FORTER FFANT FORTER FYGGGGRRINR PDTEETCHAVROGGAI  EVHENCESCHAFFART FANT FORTER FFANT FORTER FYGGGGRRINR PDTEETCHAVROGGAI  EVHENCESCHAFFART FANT FORTER FANT FORTER FANT FORTER FANT FORTER FANT FANT FANT FANT FANT FANT FANT FANT	DD170	EVVREVCSEOAETGPCRALHNRMYFDVTEGKCAPFFYGGCGGNRNNFDTEBYCMAVCGSAI	0.99	18.00	550.0	
EVURAUCSEQAETG PCRALAFRAY POVTEGKOAP PYGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PCRALET PRIF PY PYTEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PCRALET PRIF PY PYTEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PCRALET PRIF PY PYTEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PCRALET PRIF PY PYTEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PRALET PRIF PY TEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PRALET PRIF PY TEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PRALET PRIF PY TEGKOAP PY TGGGGGRINNEN PTEEY CHANGGSAI  EVURAUCSEGAETG PC PRALET PRIF PY THE PARTET PROF PARTET PARTET PROF PARTET PA	TW6176	EVVREVCSEOAETGPCRAAHFRWY FDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI			96.0	
EVVREVCEROAFTO PCRAALFRAY FDVTEGKCARP PTGGCGGRRRIN PDTEEY CHANCGSAI  EVAREVCEROAFTO FORAL FRAY FDVTEGKCAP PT GGCGGRRIN PDTEEY CHANCGSAI  EVAREVCEROAFTO FORAL FRAY FDVTEGKCAP PT GGCGGRRIN PDTEEY CHANCGSAI  EVAREVCEROAFTO FORAL FRAY FDVTEGKCAP PT GGCGGRRIN PDTEEY CHANCGSAI  EVVREVCEROAFTO FORAL FRAY FDVTEGKCAP PT GGCGGRRIN PDTEEY CHANCGSAI  EVVREVCEROAFTO FORAL FSANT FDVTEGKCAP PT GGCGGRRIN PDTEEY CHANCGSAI  EVVREVCEROAFTO FOR A FRAY FDVTEGKCAP PT GGCGGRRIN PDTEEY CHANCGSAI  EVVREVCEROAFTO FOR A FRAY FDVTEGKCAP PT GGCGGRRIN PD FEET CRANCGSAI	TW6177	EVVREVCSEQAETGPCRALHFRWY PDVTEGKCAPFFYGGCGGNRNN FDTEEYCHAVCGSAI			245.0	156.0
EVVIENCERQARIGECRALFINNY FDVTBGKCAPP FVGGCGGRRANPEDTER CCHANCGSAI  EVVIENCESQARIGECRAF FRANT FDVTEGKCAP PFVGGGGRRANR PDFESCRANCGSAI  EVVIENCESQARIGECRAF FRANT FDVTEGKCAP PFVGGGGRRANR PDFESCRANCGSAI  EVVIENCESQARIGECRAF FRANT FDVTEGKCAP PFVGGGGGRRANR PDFESCRANCGSAI  EVVIENCESQARIGEGRAF FRANT FDVTEGKCAP PFVGGGGRRANR PDFESCRANCGSAI  EVVIENCESGARIGEGRAF FRANT FDVTEGKCAP PFVGGGGRRANR PDFESCRANCGSAI  EVVIENCESCAR FRANT FOR FRANT FOR FRANT FOR FVGGGGGRRANR PFDFESCRANCGSAI  EVVIENCESCAR FRANT FOR FRANT FOR FVGGGGGRRANR PFDFESCRANCGSAI  EVVIENCESCAR FRANT FOR FVGGGGGRRANG FFDFESCRANCGSAI  EVVIENCESCAR FVGGGGGRAF FRANT FOR FVGGGGGRRANR PFDFESCRANCGSAI  EVVIENCESCAR FVGGGGGRAF FRANT FOR FVGGGGGGRRANG FFDFESCRANCGSAI  EVVIENCESCAR FVGGGGGRAF FRANT FOR FVGGGGGGRRANG FFDFESCRANCGSAI  EVVIENCESCAR FVGGGGGGRAF FRANT FFDVTEGGGGGGRRANG FFDFESCRANCGSAI  EVVIENCESCAR FVGGGGGGRAF FRANT FFDVTEGGGGGGRANG FFDFESCRANGGGGGGAI  EVVIENCESCAR FVGGGGGGRAF FFTGGGGGGGRANG FFTGGGGGGGRANG FFTGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGGG	RGOOK	EVVREVCSEOAETGPCRAALFRWYPDVTEGRCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI		12.10	0.08	
EVUREVCEROAFTO PURAL FRANT POUTBOK CADA PET GOGGORININ POTEST CHANGGSAI II  EVUREVCEROAFTO PURA FRANT POUTBOK CAP PET GOGGORININ POTEST CHANGGSAI  EVUREVCEROAFTO PURA FRANT POUTBOK CAP PET GOGGORININ POTEST CHANGGSAI  EVUREVCEROAFTO PURA FRANT POUTBOK CAP PET GOGGORININ POTEST CHANGGSAI  EVUREVCEROAFTO PURA FRANT POUTBOK PET PET GOGGORININ POTEST CHANGGSAI  EVUREVCEROAFTO PURA FRANT POUTBOK PET PET GOGGORININ POTEST CHANGGSAI  EVUREVCEROAFTO PURA FRANT POUTBOK PET PURA FRANT POTEST PURA FRANT PURA FRA	DD130	EVVREVCSEOAETGPCRALETRNYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCHAVCGSAI	5.50			9.5
EVVREVCERQAETGPCRAFFKNYFFDVTEGKCAPPTGGGGGNRNNFDTEBYCHAVGGSAI EVVREVCERQAETGPCRAFFSANYFDVTEGKCAPPTGGGGGNRNNFDTEBYCHAVGGSAI EVVREVCERQAETGPGRAALGANYFDVTEGKCAPPTGGGGSNRNNFDTEBYCHAVGGSAI EVVREVCERQAETGPGRAALGANYFDVTEGKCAPPTGGGGSNRNNFDFBYCHAVGGSAI	15100	EVVREVCSEOAETGPCRALFKRWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI		2.00	1385.0	3.3
EVVEWCERQAETGPCRAAFSANTFDVTBGKCAPFFYGGCGGNRNNFDTERCCHAVCGSAI EVVERVCERQAETGPCRAAFFEVTEGTCRAFFFYGGGGRINNIPDTERCCHAVCGSAI EVVERVCERQAETGPCRAAFFEVTEGTCRAFFFYGGGGRINNIPDTERCHAVGAICGAI	DD132	RUVREVCSEOAETGPCRAFFKRWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	=			16.8
EVVREVCSEQAETGPCRALLSAWYFDVTEGRCAPFFYGGCGGRRNNPDTEEYCHAVCGSAT	DD130	RVVREVCSEOAETGPCRAAFSAWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI				11.0
T ESCONTINUADO CONTRACTOR DE LA CONTRACTOR DEL CONTRACTOR DE LA CONTRACTOR DE LA CONTRACTOR DE LA CONTRACTOR	ונוטט	RUVREVCSEOAETGPCRALLSAWYPDVTEGRCAPFYGGCGGNRNNFDTEEYCMAVCGSAI				21.0
EVVREVCSEOAETGPCRALLWAMIFUVIEGACAFFFICGCGGMMMF DIAGETG	BG014	EVVREVCSEORETGPCRAL I WHWY FDVTEGKCAP F FYGGCGGNRNN F DTEEYCHAVCGSAI	9.20	18.70	18.0	

TOOZEGO+ OPIGOE

Title: PROTEASE INHIBITOR
PEPTIDES
Inventor(s): R. Tyler WHITE et al.
Appl. No.: 09/201,715

# FIGURE 46D

DD122	EVVREVCSEQAETGPCRALIFAMYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	15.00			46.0
BG015	EVVREVCSEQAETGPCRALIYHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	6.00	12.20	19.4	597.0
BG020	EVVREVCSEQAETGPCRAAIHKWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	1.70		106.0	
BG022	EVVREVCSEQAETGPCRAAIYHWY PDVTEGKCAPFFYGGCGGNRNN FDTEEHCMAVCGSAI	0.64	7.26	14.5	
BG023	EVVREVCSEQAETGPCRALIQHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	23.00		262.0	
BG024	EVVREVCSEQAETGPCRALIYKWYPDVTEGKCAPFFYGGCGGNRNNPDTEEYCMAVCGSAI	4.10	7.47	38.7	
BG027	EVVREVCSEORETGPCRAAIOHWYFDVTEGRCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	5.80		144.0	
DD116	EVVREVCSEOAETGPCRAAIFRWYPDVTEGKCAPFFYGGCRGNRNNFDTEEYCMAVCGSAI	0.14		583.0	84.0
FW6191	EVVREVCSEOAETGPCRAAIFRWYFDVTEGKCAPFFYGGCYGNRNNFDTEEYCMAVCGSAI	0.26		664.0	20.0
DD117	EVVREVCSEOAETGPCRALIPRMYPDVTEGKCAPFFYGGCRGNRNNFDTEEYCMAVCGSAI	0.11		1034.0	99.0
BG029	EVVREVCSEQAEVGPCRALIYHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	3.20		7.9	
BG030	EVVREVCSEQAESGPCRALIYHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	4.60		26.1	
BG033	EVVREVCSEQAEVGPCRAAIYHWYFDVTEGKCAPFFYGGCGGNRNNFDTEEHCMAVCGSAI	0.75		9.6	
BG034	EVVREVCSEOAESGPCRAAIYHWYFDVTEGKCAPFFYGGCGGRRNNFDTEEYCMAVCGSAI	0.47		18.5	
BG040	EVVREVCSEQAE IGPCRAL LYHWY PDVTEGKCAP FFYGGCGGURINIFD TEEY CMAVCGSAI	3.40		9.8	
BG016	EVVREVCSEOAETGPCRGAIQHWYFDVTEGKCAPFYGGCGGNRNNFDTEEYCMAVCGSAI	160.00		178.0	
BG017	EVVREVCSEQAETGPCRGAIRHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	180.00		200.0	
BG021	EVVREVCSEQAETGPCRGSIRHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	340.00		224.0	
BG025	EVVREVCSEQAETGPCRGLIYHWYPDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	65.00		16.2	
BG026	EVVREVCSEQAETGPCRGAIYHWYFDVTEGKCAPFFYGGCGGNRNNFDTEEYCMAVCGSAI	50.00		34.9	
DD118	EVVREVCSEQAETGPCRALHNRWYPDVTEGKCAPFFYGGCRGNRNNFDTEEYCMAVCGSAI	0.53			
DD134	EVVREVCSEQAETGPCRAL FKRWY FDVTEGKCAP FFYGGCYGNRNN FDTEEY CMAVCGSAI	1.10	1.05	15640.0	9.0
DD135	EVVREVCSEQAETGPCRALFKRMY PDVTEGKCAPFFYGGCLGNRNNFDTEEYCMAVCGSAI	1.30		7473.0	0.9
DD136	RYVVREVCSEOAETGPCBALFKBWYFDVTEGKCAPFFYGGCMGNRNNFDTEEYCMAVCGSAI	91.			~

Inventor(s): R. Tyler WHITE et al. Appl. No.: 09/201,715

Figure 47

#### VOLUMES

LES	
344.25	
245.75	
KPI	NS
298	366
266	342
354	294
258	385
168	288
266	469
172	338
184	272
245.75	344.25
66.2414415	63.97488346
-	0.009094999
	344.25 245.75 KPI 298 266 354 258 168 266 172 184 245.75

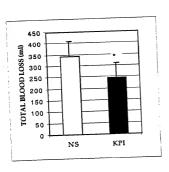


Figure 48

#### HEMOGLOBIN

NS	23	1.61
KPI	13	.59
	KPI	NS
	16	5.58
	15	5.19

71	No	
	16.58	24.95
	15.19	24.87
	20.21	20.46
	8.99	27.59
	14.63	18.23
	15.31	31.59
	7.7	23.26
	10.14	17.96
2	E007E 00	61275

MEAN 13.59375 23.61375 STDEV 4.261438 4.68761 TTEST 0.000536

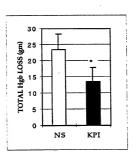


Figure 49

PaO2

	Baseline	PaO2		End CPB	
	KPI	NS		KPI	NS
	652.2	670.9	ŀ	495.7	60.5
	654	559.2		444.6	132.2
	596.2	622.9		170.2	93.8
	606.2	689.2		264.2	333.9
	633.1	665.1		567.2	341.7
	646.6	527		507.4	226.9
	563.2	461.7		547.1	89.1
	659.9	508		416.6	59.7
MEAN	626.425	588		426.625	167.225
STDEV	34.4692	85.5055		140.474	117.993
	3	6		1	1
TTEST	p=	0.268 N.S.		p=	0.0014
		-			

Obs 60 n	nin	
KPI	NS	
483.7	441.3	
330.1	448.7	
415.4	85.1	
430.2	529.6	
613	568.3	
564.3	438.1	
501	42.6	
504.5	405.8	
480.275	369.938	
88.6187	196.523	
9	5	
=מ	0.17915	

Obs 180	min
KPI	NS
	391.3
264.1	484.6
416.5	81.3
361.9	333.2
90.8	546.6
518.2	485.3
494.2	45.6
452	383.7
371.1	344
150.277	186.22
4	. 7
p=	0.76

Figure 50

Summary of Data

	min	6.2	2.7	7.1	1.9	6.7	6.3	4.4	4.2	5.3	1.72	 5.4	7.1	6.5	6.3	9.6	7.4	4.2	5.3	6.1	0.85	
	120-180min																					1
540	0-30min 30-60min 60-120min	8.6	6.7	7	4.4	7	5.6	5.4	5.4	6.26	1.32	6.1	9.7	7.5	7.1	5.2	7.3	5.8	9	6.58	16:0	
st tube Hb	30-60min	4.3	6.4	4.4	4	6.5	6.1	4.6	5.8	5.26	1.04	8.6	7.4	7.5	7.2	7.2	7	7.7	8.2	7.6	1.04	
Serial Chest tube Hbg	0-30min	3.7	4.3	4.1	2.8	6.3	4.1	3.1	6.9	4.41	1.45	7.7	7.2	5.4	8.4	7.5	4	7.5	7.4	689	1.44	
										MEAN	STDEV							_		MEAN	STDEV	
nes	Sacrifice	113	89	212	89	72	28	38	56			92	106	42	82	148	208	120	99			
Total Volumes	Chest tube Sacrifice	185	198	142	190	8	188	134	158			274	236	252	303	140	261	218	506			
Tc	Total Hgb Loss	16.58	15.19	20.21	8.99	14.63	15.31	7.7	10.14	13.59	4.26	24.95	24.87	20.46	27.59	18.23	31.59	23.26	17.96	73.61	4 69	, C.F.
	Total volume loss	298	266	354	258	168	266	172	184	245.75	66.24	366	342	294	385	288	469	338	272	244.75	20 63	72.50
		KPI-1	KPI-2	KPI-3	KPI-4	KPI-5	KPI-6	KPI-7	KPI-8	MEAN	STDEV	NS-1A	NS-2	NS-3	NS-4	NS-5	9-SN	NS-7	NS-8	N. E. W.	VICTORY	SIUEV